

# In-Class Lab 9

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The purpose of this lab is to practice using R to test for serial correlation and how to fix it. The lab may be completed as a group. To receive credit, upload your .R script to the appropriate place on eCampus (“In-Class Labs” folder).

## For starters

Open a new R script (named ICL9\_XYZ.R, where XYZ are your initials)

## Clean out/“Sweep’’ R Studio

Click the broom in the Environment panel (top-right), it is directly below the Tutorial button. Also, in the bottom-right panel, click the Plots button and then click the broom in that panel. This should help with loading things into R.

## R Packages

First, install the `pdfetch`, `tsibble`, and `COVID19` packages. `pdfetch` stands for “Public Data Fetch” and is a slick way of downloading statistics on stock prices, GDP, inflation, unemployment, etc. `tsibble` is a package useful for working with time series data. `COVID19` pulls up-to date data on COVID-19 cases, deaths, etc.

```
install.packages("COVID19")
install.packages("sandwich")
install.packages("lmtest")
install.packages("pdfetch")
install.packages("tsibble")
install.packages("magrittr")
install.packages("tidyverse")
```

```
library(wooldridge)
library(car)
library(magrittr)
library(lmtest)
library(pdfetch)
library(sandwich)
library(tsibble)
library(COVID19)
library(tidyverse)
```

## Load the data

We're going to use data on US COVID-19 cases, death, and other information

```
COVID_Data <- covid19(c("US"))
COVID_Time_Series <- as_tsibble(COVID_Data, key = id, index = date)
```

Now it will be easy to include lags of various variables into our regression models

```
View(COVID_Time_Series)
```

## Plot time series data

Let's have a look at the data on new daily cases and deaths:

```
COVID_Time_Series %>% mutate(new_deaths = difference(deaths), # data comes in cumulative format
                                new_tests = difference(tests), # "difference" converts it to
                                new_cases = difference(confirmed)) # "new cases" etc.

# plots
ggplot(COVID_Time_Series, aes(date, new_cases)) + geom_line()

## Warning: Removed 19 rows containing missing values ('geom_line()').

ggplot(COVID_Time_Series, aes(date, new_deaths)) + geom_line()

## Warning: Removed 19 rows containing missing values ('geom_line()').

# plots with 7-day rolling average
ggplot(COVID_Time_Series, aes(date, new_cases)) + geom_line(aes(y=rollmean(new_cases, 7,
                                na.pad=TRUE)))

## Warning: Removed 25 rows containing missing values ('geom_line()').

ggplot(COVID_Time_Series, aes(date, new_deaths)) + geom_line(aes(y=rollmean(new_deaths, 7,
                                na.pad=TRUE)))

## Warning: Removed 25 rows containing missing values ('geom_line()').
```

## Determinants of US COVID-19 Cases

Now let's estimate the following regression model:

$$\log(new\_cases_t) = \beta_0 + \beta_1 gath_t + \beta_2 gath_{t-7} + \beta_3 gath_{t-14} + \beta_4 \log(new\_cases_{t-7}) + u_t$$

where *new\_cases* is the number of new COVID cases, and *gath* is a variable taking on values 0–4 representing severity of gatherings restrictions.

```

COVID_Time_Series %>% mutate(log.new.cases = log(new_cases),
                                log.new.cases = replace(log.new.cases,new_cases==0,NA_real_))

## Warning: There was 1 warning in 'mutate()'.
## i In argument: 'log.new.cases = log(new_cases)'.
## Caused by warning in 'log()':
## ! NaNs produced

regression <- lm(log.new.cases ~ gatherings_restrictions + lag(gatherings_restrictions,7) +
                    lag(gatherings_restrictions,14) + lag(log.new.cases,7),
                    data=COVID_Time_Series)
summary(regression)

##
## Call:
## lm(formula = log.new.cases ~ gatherings_restrictions + lag(gatherings_restrictions,
##                 7) + lag(gatherings_restrictions, 14) + lag(log.new.cases,
##                 7), data = COVID_Time_Series)
##
## Residuals:
##       Min     1Q   Median     3Q    Max 
## -1.87003 -0.18663 -0.02246  0.18695  1.96682
##
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)    
## (Intercept)             1.400710  0.082420 16.995 < 2e-16 ***
## gatherings_restrictions -0.262853  0.035866 -7.329 4.68e-13 ***
## lag(gatherings_restrictions, 7)  0.012143  0.053460  0.227    0.82  
## lag(gatherings_restrictions, 14)  0.255587  0.035775  7.144 1.71e-12 ***
## lag(log.new.cases, 7)        0.876298  0.007888 111.097 < 2e-16 ***
## ---                        
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.4115 on 1033 degrees of freedom
##   (138 observations deleted due to missingness)
## Multiple R-squared:  0.9286, Adjusted R-squared:  0.9283 
## F-statistic: 3357 on 4 and 1033 DF,  p-value: < 2.2e-16

```

## Testing for Serial Correlation

Using the Durbin-Watson Test:

```
durbinWatsonTest(regression)
```

```
##   lag Autocorrelation D-W Statistic p-value
##   1      0.4509892     1.086493      0
## Alternative hypothesis: rho != 0
```

Now using the Lagrange Multiplier Test:

```

bgtest(regression)

##
## Breusch-Godfrey test for serial correlation of order up to 1
##
## data: regression
## LM test = 211.85, df = 1, p-value < 2.2e-16

```

## Correcting for Serial Correlation

Now let's compute Newey-West standard errors. To do so, we'll use the `vcov` option in the `modelsummary()` function along with the `NeweyWest` function from the `sandwich` package.

```

NW_VCOV <- NeweyWest(regression)
coeftest(regression, vcov = NW_VCOV)

```

```

##
## t test of coefficients:
##
##                               Estimate Std. Error t value Pr(>|t|)
## (Intercept)                 1.400710  0.323312 4.3324 1.619e-05 ***
## gatherings_restrictions    -0.262853  0.104689 -2.5108  0.01220 *
## lag(gatherings_restrictions, 7) 0.012143  0.085269  0.1424  0.88679
## lag(gatherings_restrictions, 14) 0.255587  0.106085  2.4093  0.01616 *
## lag(log.new.cases, 7)        0.876298  0.028811 30.4150 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

How does your interpretation of the effect of gathering restrictions change after using the Newey-West standard errors?